

Supplemental Material.

Norton et al, Circulation Cardiovascular Genetics 2013: *Exome sequencing and genome-wide linkage analysis in 17 families illustrates the complex contribution of TTN truncating variants to dilated cardiomyopathy.*

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HeartGO:

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NHLBI GO Exome Sequencing Project

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¹⁰Lung Project Team, ¹¹Personal Genomics Project Team, ¹²Phenotype and Harmonization Working Group, ¹³Population Genetics and Statistical Analysis Working Group, ¹⁴Publications and Presentations Working Group, ¹⁵Quantitative Analysis Ad Hoc Task Group, ¹⁶Sequencing and Genotyping Working Group, ¹⁷Steering Committee, ¹⁸Stroke Project Team, ¹⁹Structural Variation Working Group, ²⁰Subclinical/Quantitative Project Team

ESP Cohorts

²¹Acute Lung Injury (ALI), ²²Atherosclerosis Risk in Communities (ARIC), ²³Cardiovascular Health Study (CHS), ²⁴Chronic Obstructive Pulmonary Disease (COPD) Gene, ²⁵Coronary Artery Risk Development in Young Adults (CARDIA), ²⁶Cystic Fibrosis (CF), ²⁷Early Pseudomonas Infection Control (EPIC), ²⁸Framingham Heart Study (FHS), ²⁹Jackson Heart Study (JHS), ³⁰Lung Health Study (LHS), ³¹Multi-Ethnic Study of Atherosclerosis (MESA), ³²Pulmonary Arterial Hypertension (PAH), ³³Severe Asthma Research Program (SARP), ³⁴Women's Health Initiative (WHI)